

FIG. 1A



FIG. 1B

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Fig. 2a



Fig. 2b

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<u>HRS.</u>		N	STANDARD DEVIATION	STANDARD ERROR
0	100.000%	12	0.000%	0.000%
24	100.000%	12	0.000 <i>%</i>	0.000%
48	96.740%	12	5.107%	1.474%
72				
96	NA		######################################	
120	90.218%	12	3.763%	1.086%
144	NA	*****		
168	88.768 %	12	2.235%	0.645%
192	84.058%	12	5.664%	1.635%
216	80.435%	12	6.272%	1.811%
240	73.553%	12	9.536%	2.753%
264	NA			144001400700000000000000000000000000000
288	67.391%	12	8.987%	2.594%
MINI-	PATCHES (HSI	7-12):		
HRS.	AVERAGE	N	STANDARD DEVIATION	STANDARD ERROR
0	100.000%	4	0.000%	0.000%
24	100.000%	4	0.000%	0.000%
48	100.000%	4	0.000%	0.000%
72	NA	*		
96	NA	****		
120	94.563%	4	2.175%	1.088%
144	NA		\	
168	93.475%	4	2.511%	1.256%
192	93.475%	4	2.511%	1.256%
216	93.475%	4	2.511%	1.256%
240	93.475%	4	2.511%	1.256%
264	NA			
288	92.388%	4	2.175%	1.088%
MINI-	PATCHES (F-12	2):		
HRS.	AVERAGE	N	STANDARD DEVIATION	STANDARD ERROR
0	100.000%	12	0.000%	0.000%
24	100.000%	12	0.000%	0.000%
48	100.000%	12	0.000%	0.000%
72	NA			,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,
96	NA			
120	99.638%	12	1.256%	0.363%
144	NA			
168	99.638%	12	1.256%	0.363%
192	99.638%	12	1.256%	0.363%
216	99.638%	12	1.256%	0.363%
240	99.638%	12	1.256%	0.363%
3.7.4	NA			
264	1427			
288 288	99.638%	12	1.256%	0.363%

FIG. 3A TABLE 1

FIGURE 3B

20ml- Full	Size PAP	(HSDMEM)
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Hrs.	Average	Standard D	viation Standard Error
0	100.000 %	± 0.000%	± 0.000 %
24	100.000%	± 0.000%	± 0.000%
48	100.000%	± 0.000%	± 0.000%
72	99.690%	± 1.037%	± 0.299%
96	91.350%	± 4.812%	± 1.389%
120	NA	NA	
144	83.330%	± 7.693%	± 2.221
168	61.770%	± 9.177	± 2.649

FIG. 4A TABLE 2

Matrix Reorganization PAP

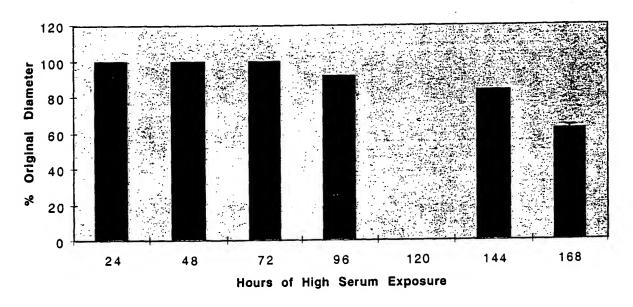


FIG. 4B. GRAPH 2



FIG. 5A



FIG. 5B

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FIG. 6A



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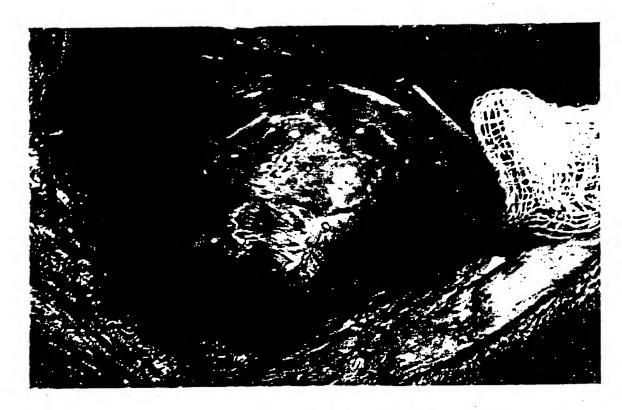


FIG. 7A

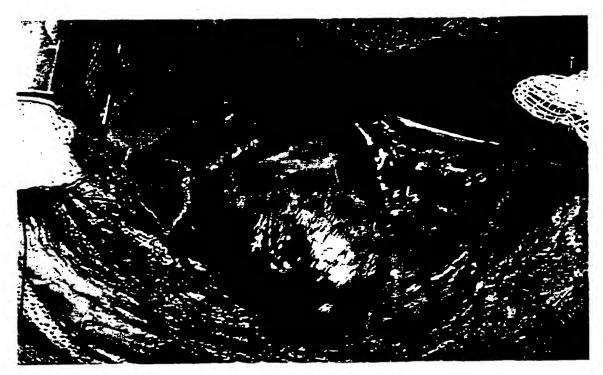


FIG. 7B

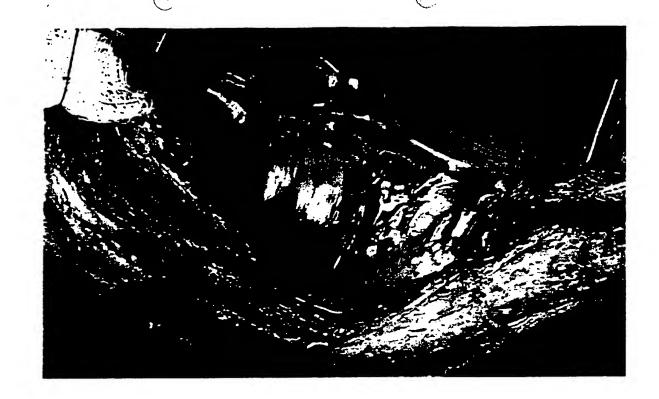


FIG. 7C

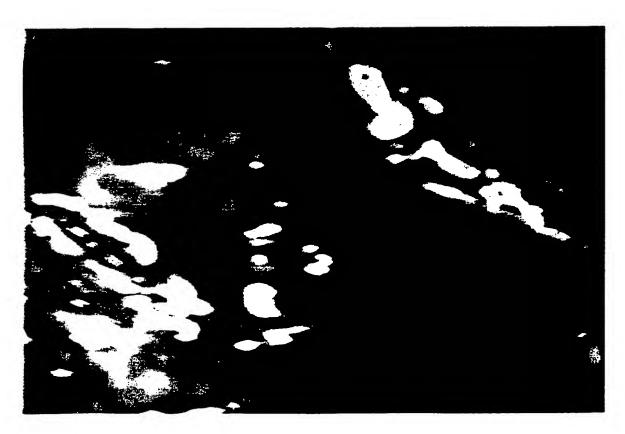


FIG. 7D



FIG. 8A



FIG. 8B



FIG. 8C



FIG. 8D

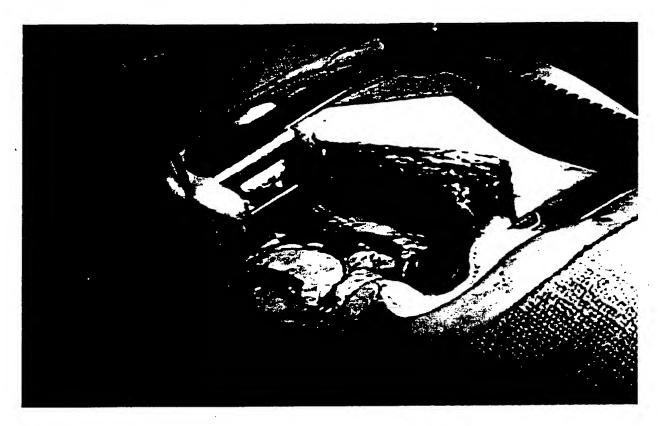


FIG. 8E

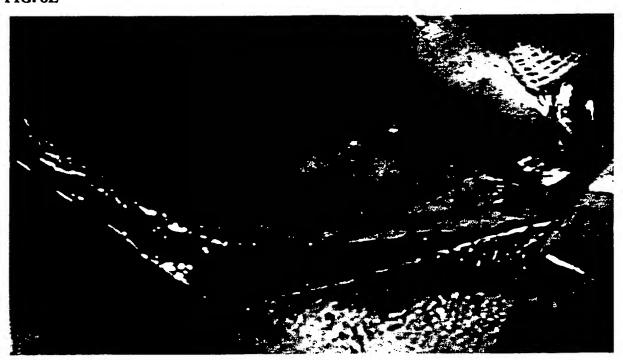


FIG. 9A



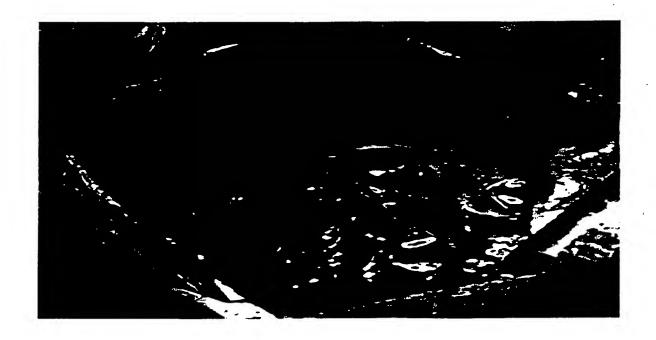


FIG. 9C



FIG. 9D

